EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	0	pyruvate adj decarboxylase and etanol and yeast and gene and bacillus	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L2	0	pyruvate adj decarboxylase and etanol	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L3	3478	pyruvate adj decarboxylase and ethanol	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:23
L4	1073	13 and gene and bacillus	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:25
L5	924	13 and gene and bacillus and alcohol adj dehydrogenase	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:26
L6	431	I5 and gram-positive	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:27
L7	29	I5 and gram-positive and lactate adj dehydrogenase	US-PGPUB; USPAT; DERWENT	OR	ON	2006/08/04 16:27

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NEWS 14 JUl 14 FSTA enhanced with Japanese patents

NEWS 15 JUl 19 Coverage of Research Disclosure reinstated in DWPI

NEWS EXPRESS JUNE 30 CURRENT WINDOWS VERSION IS V8.01b, CURRENT MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP), AND CURRENT DISCOVER FILE IS DATED 26 JUNE 2006.

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=> s ethanol and yeast and gene
L1 3636 ETHANOL AND YEAST AND GENE

=> s l1 and bacillus

L2 90 L1 AND BACILLUS

=> s 12 and pyruvate (w) decarboxylase 7 L2 AND PYRUVATE (W) DECARBOXYLASE L3

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ANSWER 1 OF 7 MEDLINE on STN L3

ACCESSION NUMBER: 2005658391 MEDLINE

DOCUMENT NUMBER:

PubMed ID: 16339947

TITLE:

Construction and expression of an ethanol production operon in Gram-positive bacteria.

AUTHOR:

Talarico Lee A; Gil Malgorzata A; Yomano Lorraine P; Ingram

Lonnie O; Maupin-Furlow Julie A

CORPORATE SOURCE:

Department of Microbiology and Cell Science, University of

Florida, Gainesville, FL 32611-0700, USA.

SOURCE:

Microbiology (Reading, England), (2005 Dec) Vol. 151, No.

Pt 12, pp. 4023-31.

Journal code: 9430468. ISSN: 1350-0872.

PUB. COUNTRY:

England: United Kingdom

DOCUMENT TYPE:

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

200602

ENTRY DATE:

Entered STN: 18 Dec 2005

Last Updated on STN: 1 Mar 2006 Entered Medline: 28 Feb 2006

AB Pyruvate decarboxylase (PDC), an enzyme central to homoethanol fermentation, catalyses the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host Bacillus megaterium. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive Sarcina ventriculi was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of Geobacillus stearothermophilus. The resulting Gram-positive ethanol production operon was expressed at high levels in B. megaterium. Extracts from this recombinant were shown to catalyse the production of ethanol from pyruvate.

ANSWER 2 OF 7 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN L3

ACCESSION NUMBER: DOCUMENT NUMBER:

2006:151966 BIOSIS PREV200600152190

TITLE:

Construction and expression of an ethanol production operon in Gram-positive bacteria.

AUTHOR (S):

Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine P.; Ingram, Lonnie O.; Maupin-Furlow, Julie A. [Reprint Author]

CORPORATE SOURCE:

Univ Florida, Dept Microbiol and Cell Sci, Gainesville, FL

32611 USA

jmaupin@ufl.edu

SOURCE:

Microbiology (Reading), (DEC 2005) Vol. 151, No. Part 12,

pp. 4023-4031. ISSN: 1350-0872.

DOCUMENT TYPE:

Article

LANGUAGE:

English

ENTRY DATE:

Entered STN: 1 Mar 2006

Last Updated on STN: 1 Mar 2006

AB Pyruvate decarboxylase (PDC), an enzyme central to

homoethanol fermentation, catalyses the non-oxidative decarboxylation of

pyruvate to acetaldehyde with release of carbon dioxide. 1 enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host Bacillus megaterium. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive Sarcina ventriculi was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of Geobacillus stearothermophilus. The resulting Gram-positive ethanol production operon was expressed at high levels in B. megaterium. Extracts from this recombinant were shown to catalyse the production of ethanol from pyruvate.

L3 ANSWER 3 OF 7 CAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 2006:17838 CAPLUS

DOCUMENT NUMBER: 144:405166

TITLE: Construction and expression of an ethanol production operon in Gram-positive bacteria

AUTHOR(S): Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine

P.; Ingram, Lonnie O.; Maupin-Furlow, Julie A.

CORPORATE SOURCE: Department of Microbiology and Cell Science,

University of Florida, Gainesville, FL, 32611-0700,

USA

SOURCE: Microbiology (Reading, United Kingdom) (2005),

151(12), 4023-4031

CODEN: MROBEO; ISSN: 1350-0872 Society for General Microbiology

DOCUMENT TYPE: Journal LANGUAGE: English

PUBLISHER:

Pyruvate decarboxylase (PDC), an enzyme central to homoethanol fermentation, catalyzes the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-pos. biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-pos. host Bacillus megaterium. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this anal., the pdc gene of Gram-pos. Sarcina ventriculi was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-pos. This gene was thus selected for transcriptional coupling to the alc. dehydrogenase gene (adh) of Geobacillus stearothermophilus. The resulting Gram-pos. ethanol production operon was expressed at high levels in B. megaterium. Exts. from this recombinant were shown to catalyze the production of ethanol from pyruvate.

REFERENCE COUNT: 35 THERE ARE 35 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L3 ANSWER 4 OF 7 CAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 1992:649986 CAPLUS

DOCUMENT NUMBER: 117:249986

TITLE: Ethanol production by by bacteria carrying

foreign genes for alcohol dehydrogenase and

pyruvate decarboxylase

INVENTOR(S): Ingram, Lonnie O.; Beall, David S.; Burchhardt,

Gerhard F. H.; Guimaraes, Walter V.; Ohta, Kazuyoshi;

Wood, Brent E.; Shanmugam, Keelnatham T.; Fowler,

David A.; Ben-Bassat, Arie

PATENT ASSIGNEE(S): University of Florida, USA; Bioenergy International,

L.C.

SOURCE: PCT Int. Appl., 153 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

10

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PA	PATENT NO.					KIND DATE			AP	APPLICATION NO.				DATE			
WO	9216615			A1 19921001			WO 1992-US1807			19920318							
	W:	ΑT,	AU,	BB,	BG,	BR,	CA,	·CH,	CS, DI	E, DK,	ES,	FI,	GB,	HU	I, JP,	ΚP,	
		KR,	LK,	LU,	MG,	MN,	MW,	NL,	NO, P	L, RO,	RU,	SD,	SE,	US	;		
	RW:	AT,	BE,	BF,	ВJ,	CF,	CG,	CH,	CI, CI	M, DE,	DK,	ES,	FR,	GΑ	GB,	GN,	
		GR	TT	T.IJ.	MC .	MT.	MR.	NT.	SE. SI	N. TD.	TG						
US	5424	202			Α		1995	0613	US	1992-	-8463	44			19920	306	
AU	9217	794			AI		1992	T02T	AU	1992.	- 1 / / 9	4			19920	318	
AU	6727	48			В2		1996	1017	CN								
CN	1070	424			A		1993	0331	CN	1992-	-1018	77			19920	318	
CN	1065	915			В		2001	0516									
EP	5766	21			A1		1994	0105	EP	1992-	-9109	33			19920	318	
	5766																
	R:	ΑT,	BE,	CH,	DE,	DK,	ES,	FR,	GB, G	R, IT,	LI,	LU,	MC,	NL	, SE		
JР	0650	5875			T2		1994	0707	JP	1992-	-5099	41			19920	318	
JP	3457	664			B2		2003	1020	BR								
BR	9205	782			Α		1994	0726	BR	1992	-5782				19920	318	
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NO	9303	178			Α		1993	1108	AT NO	1993	-3178				19930	907	
NO	3155	67			В1		2003	0922									
CN	1342	773			A		2002	0403	CN	2000-	-1317	79			20001	020	
AU	2005	24892	24		A1		2006	0202	UA	2005	-2489	24			20051	223	
PRIORIT	Y APP	LN.	INFO	. :					US	1991	-6708	21		Α	19910	318	
									US	1992	-8463	44		Α	19920	306	
									US	1988	-2390	99		B2	19880	831	
									US	1989	-3520	62		A2	19890	515	
									US US US US WO	1990	-6242	77		B2	19901	207	
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Bacterial hosts, excluding Escherichia coli, expressing heterologous AB genes for alc. dehydrogenase (I) and pyruvate decarboxylase (II) are used for manufacture of EtOH. II is used to prevent accumulation of acid metabolites. Plasmids, e.g. pLOI555 carrying genes for I and II of Zymomonas mobilis driven by the lac promoter, are provided for preparation of the host. The method is further improved by transforming the host with genes for proteins that facilitate transport and metabolism of oligosaccharides, e.g., of C5-6 sugars, which host is, preferably, also expressing a heterologous gene for a polysaccharase such as a cellulolytic enzyme, a xylanolytic enzyme, or a starch-degrading enzyme. These hosts also preferably express heterologous genes for polysaccharide- degrading enzymes (e.g. those degrading cellulose, xylans, or starch). A cost-effective fermentation process for manufacturing EtOH from oligosaccharide feedstocks using a single, genetically engineered microorganism is also disclosed. An ethanologenic strain Klebsiella oxytoca M5A1(pLOI555) was prepared and was further transformed with plasmid pLOI2003 encoding xylanase (gene xynZ) and xylosidase (gene xylB) of Clostridium thermocellum to obtain a transformant capable of converting xylan to EtOH.

L3 ANSWER 5 OF 7 LIFESCI COPYRIGHT 2006 CSA on STN

ACCESSION NUMBER: 2006:26910 LIFESCI

TITLE: Construction and expression of an ethanol

production operon in Gram-positive bacteria

AUTHOR: Talarico, Lee A.; Gil, Malgorzata A.; Yomano, Lorraine P.;

Ingram, Lonnie O.; Maupin-Furlow, Julie A.

CORPORATE SOURCE: Department of Microbiology and Cell Science, University of

Florida, Gainesville, FL 32611-0700, USA; E-mail:

jmaupin@ufl.edu

SOURCE: Microbiology, (20051200) vol. 151, no. 12, pp. 4023-4031.

ISSN: 1350-0872.

DOCUMENT TYPE: Journal FILE SEGMENT: W2; J LANGUAGE: English

SUMMARY LANGUAGE: English

Pyruvate decarboxylase (PDC), an enzyme central to

homoethanol fermentation, catalyses the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host Bacillus megaterium. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive Sarcina ventriculi was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of Geobacillus stearothermophilus. The resulting Gram-positive ethanol production operon was expressed at high levels in B. megaterium. Extracts from this recombinant were shown to catalyse the production of ethanol from pyruvate.

L3 ANSWER 6 OF 7 LIFESCI COPYRIGHT 2006 CSA on STN

ACCESSION NUMBER:

84:75038 LIFESCI

TITLE:

.

Xylose-isomerase in yeast.

AUTHOR:

Wilhelm, M.; Erhart, E.; Hollenberg, C.P.

CORPORATE SOURCE:

Inst. Mikrobiol., Univ. Duesseldorf, Duesseldorf, FRG

SOURCE:

RIV. BIOL., (1984) vol. 77, no. 4, pp. 607-608.

Meeting Info.: International Course on Microbial Breeding.

Spoleto (Italy). 3-8 Sep 1984.

DOCUMENT TYPE:

Journal TREATMENT CODE: Conference FILE SEGMENT: G; W; N; K LANGUAGE: English

Saccharomyces cerevisiae is not able to ferment xylose to ethanol . To explore the feasibility of constructing yeast strains that can produce ethanol from xylose, the authors have decided to introduce a xylose isomerase gene into S. cerevisiae under control of the yeast pyruvate decarboxylase promoter and to study the effects of its expression. As a first step the authors have isolated a BamHi fragment from Bacillus subtilis which carries the genetic information for xylose isomerase and exlyulokinase.

L3 ANSWER 7 OF 7 EMBASE COPYRIGHT (c) 2006 Elsevier B.V. All rights reserved on STN

ACCESSION NUMBER:

2005583341 EMBASE

TITLE:

Construction and expression of an ethanol production operon in Gram-positive bacteria.

AUTHOR:

Talarico L.A.; Gil M.A.; Yomano L.P.; Ingram L.O.;

Maupin-Furlow J.A.

CORPORATE SOURCE:

J.A. Maupin-Furlow, Department of Microbiology and Cell Science, University of Florida, Gainesville, FL 32611-0700,

United States. jmaupin@ufl.edu

SOURCE: Microbiology, (2005) Vol. 151, No. 12, pp. 4023-4031. .

Refs: 35

ISSN: 1350-0872 CODEN: MROBEO

COUNTRY: United Kingdom
DOCUMENT TYPE: Journal; Article
FILE SEGMENT: 004 Microbiology

LANGUAGE: English SUMMARY LANGUAGE: English

ENTRY DATE: Entered STN: 12 Jan 2006

Last Updated on STN: 12 Jan 2006 Pyruvate decarboxylase (PDC), an enzyme central to homoethanol fermentation, catalyses the non-oxidative decarboxylation of pyruvate to acetaldehyde with release of carbon dioxide. PDC enzymes from diverse organisms have different kinetic properties, thermal stability and codon usage that are likely to offer unique advantages for the development of desirable Gram-positive biocatalysts for use in the ethanol industry. To examine this further, pdc genes from bacteria to yeast were expressed in the Gram-positive host Bacillus megaterium. The PDC activity and protein levels were determined for each strain. In addition, the levels of pdc-specific mRNA transcripts and stability of recombinant proteins were assessed. From this analysis, the pdc gene of Gram-positive Sarcina ventriculi was found to be the most advantageous for engineering high-level synthesis of PDC in a Gram-positive host. This gene was thus selected for transcriptional coupling to the alcohol dehydrogenase gene (adh) of Geobacillus stearothermophilus. The resulting Gram-positive ethanol production operon was expressed at high levels in B. megaterium. Extracts from this recombinant were shown to catalyse the

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production of ethanol from pyruvate. .COPYRGT. 2005 SGM.

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